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APPLICATION NO.	FILING DATE	ILING DATE FIRST NAMED INVENTOR		CONFIRMATION NO.	
10/573,332	04/06/2007	David W. Morris	PP023362.0003	5041	
	7590 03/03/201 ACCINES AND DIAG	EXAMINER			
INTELLECTUA	AL PROPERTY- X10	HARRIS, ALANA M			
P.O. BOX 8097 Emeryville, CA			ART UNIT PAPER NUMBER		
•			1643		
			MAIL DATE	DELIVERY MODE	
			03/03/2010	PAPER	

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

	Application No.	Applicant(s)
	10/573,332	MORRIS ET AL.
Office Action Summary	Examiner	Art Unit
	Alana M. Harris, Ph.D.	1643
The MAILING DATE of this communication app Period for Reply	ears on the cover sheet with the c	orrespondence address
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication If NO period for reply is specified above, the maximum statutory period v - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be time will apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	lely filed the mailing date of this communication. (35 U.S.C. § 133).
Status		
1)☒ Responsive to communication(s) filed on 17 No. 2a)☐ This action is FINAL. 2b)☒ This 3)☐ Since this application is in condition for alloware closed in accordance with the practice under Example 2.	action is non-final. nce except for formal matters, pro	
Disposition of Claims		
 4) Claim(s) 1-55 is/are pending in the application. 4a) Of the above claim(s) 1-32 and 37-55 is/are 5) Claim(s) is/are allowed. 6) Claim(s) 33-36 is/are rejected. 7) Claim(s) is/are objected to. 8) Claim(s) are subject to restriction and/or 	e withdrawn from consideration.	
Application Papers		
9) The specification is objected to by the Examine 10) The drawing(s) filed on is/are: a) acce Applicant may not request that any objection to the examine Replacement drawing sheet(s) including the correct 11) The oath or declaration is objected to by the Examine	epted or b) objected to by the Eddrawing(s) be held in abeyance. See ion is required if the drawing(s) is obj	e 37 CFR 1.85(a). ected to. See 37 CFR 1.121(d).
Priority under 35 U.S.C. § 119		
 12) Acknowledgment is made of a claim for foreign a) All b) Some * c) None of: 1. Certified copies of the priority documents 2. Certified copies of the priority documents 3. Copies of the certified copies of the prior application from the International Bureau * See the attached detailed Office action for a list 	s have been received. s have been received in Application rity documents have been receive u (PCT Rule 17.2(a)).	on No ed in this National Stage
Attachment(s)	A> □ Indoor : 0	(DTO 442)
 Notice of References Cited (PTO-892) Notice of Draftsperson's Patent Drawing Review (PTO-948) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date <u>04/10/2008</u>. 	4) Interview Summary Paper No(s)/Mail Da 5) Notice of Informal P 6) Other:	ite

Art Unit: 1643

DETAILED ACTION

Election/Restrictions

1. Applicant's election with traverse of Group VIII (claims 33 and 34) in the reply filed on November 17, 2010 is acknowledged. The traversal is on the ground(s) that "Groups VIII and IX (claims 35 and 36) are both directed to embodiments that comprise at least two polynucleotides" and consequently a search of both sets of claims would be duplicative and "...should not be an undue burden on the Examiner". This is found persuasive and both Groups will be examined.

The remainder of the requirement is still deemed proper and is therefore made FINAL.

2. Claims 1-55 are pending.

Claims 1-32 and 37-55, drawn to non-elected inventions are not examined on the merits.

Claims 33-36 are examined on the merits.

Art Unit: 1643

Claim Rejections - 35 USC § 112

3. The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

- 4. Claims 35 and 36 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.
- a. Claims 35 and 36 are drawn to an electronic library. It is not clear if Applicants intended to state an electronic medium, an array or microarray that comprises sequences. The term, electronic library has not been defined in the specification. Accordingly, the metes and bounds cannot be determined.

Claim Rejections - 35 USC § 102

5. The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351(a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

(e) the invention was described in a patent granted on an application for patent by another filed in the United States before the invention thereof by the applicant for patent, or on an international application by another who has fulfilled the requirements of paragraphs (1), (2), and (4) of section 371(c) of this title before the invention thereof by the applicant for patent.

Application/Control Number: 10/573,332

Page 4

Art Unit: 1643

The changes made to 35 U.S.C. 102(e) by the American Inventors

Protection Act of 1999 (AIPA) and the Intellectual Property and High Technology

Technical Amendments Act of 2002 do not apply when the reference is a U.S.

patent resulting directly or indirectly from an international application filed

before November 29, 2000. Therefore, the prior art date of the reference is

determined under 35 U.S.C. 102(e) prior to the amendment by the AIPA (pre
AIPA 35 U.S.C. 102(e)).

6. Claims 33 and 35 are rejected under 35 U.S.C. 102(e) as being anticipated by Knoll et al./ U.S. Patent number 7,014,997 B2 (filed May 14, 2001). Knoll discloses sequences that would selectively hybridize to Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, fragments thereof or their complements within in a sample, see sequence alignment information following instant rejection; column 1, lines 36-45; and column 14, lines 56-65. Knoll discloses sequences 425 and 148, which are the same as Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, respectively, corresponding hybridization probes, arrays, and kits in which these components are contained, see abstract; column 2, lines 5-50; column 3, lines 17-25; bridging paragraph of columns 24 and 25; and the claims beginning in column 37. These components are useful in detecting "[c]hromosomal abnormalities often common and ...diagnostic in...leukemia and other cancers", see column 2, 1st full sentence.

In absence of a clear definition of the term, electronic library as cited in the pending 112, 2nd paragraph rejection, the art reads on claim 35. The Examiner regards the cited arrays as the equivalent of Applicants' electronic library.

Search results between Applicants' SEQ ID NO: 4 and Knoll sequence 425.

```
US-09-854-867-425, rni
; Sequence 425, Application US/09854867
; Patent No. 7014997
; GENERAL INFORMATION:
 APPLICANT: JOAN, KNOLL H
; APPLICANT: ROGAN, PETER K
; TITLE OF INVENTION: SINGLE COPY GENOMIC HYBRIDIZATION PROBES AND METHOD OF GENERATING SAME
; FILE REFERENCE: 30307
  CURRENT APPLICATION NUMBER: US/09/854,867
 CURRENT FILING DATE: 2003-05-08
; NUMBER OF SEQ ID NOS: 613
  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 425
  LENGTH: 3285
  TYPE: DNA
   ORGANISM: Homo sapiens
  FEATURE:
 Query Match
                    24.1%; Score 289.6; DB 5; Length 3285;
 Best Local Similarity 65.9%;
 Matches 567; Conservative 5; Mismatches 216; Indels 72; Gaps
        34 CAAAGAAATAAGAGAAACACATAAAGCAAGAAATAAGAGAAACACATAATTCAACAATAA 93
           365 CAAAATACATAAAGCAAATACTAATAGAACTGAAAAGAGAAATAGACAAATCCACAATAA 424
Qу
        94 TA----GAAACTTAAATATCCTCCTTTCAATAATAGATACAACAACTAAGCAGTTGATCA 149
           425 TAGTTGGAGACTTCAATACCCCACTGTCAGTAATTGATAGATCAACTAGACAGAAAATCA 484
        150 ACAAGAAAACAGAAGATTTGAACAATGCTATGAACCAACTAGACCTAACGTCTA---TCT 206
           485\ ATAAGGATATAGAAGACTTGAACAACACTATCAACCAACTGGACCTAATTGACATATTAT\ 544
Db
        207 ATAAAACACCACCCAACAACAGCAGAATACATATTCTTCTCAGATATACATAGAACATTC 266
Qу
           545 AGAACACTCCACCCAACAACAGCAGAATACACATTCTTYTCAAGTGCACATGGAACATTC 604
Db
        267 TCCAGGATAGGCCATCTGTTAGGACATAAAACAAGTCTCAAAAAATGTAAAAGAATTGAG 326
Q.y
            605 ACCAAGATAGACCATATGCTRGGCCATAAAACAAGTCTCAATAAATTTAAAAAAAATTGAA 664
        327 ATCAGACAAAGTCTGTTCTCTGACCACAA-------CCAGTAACAGAAG 368
QУ
           665 ATYATACAAAGTATGTTCTCTGACCACAATGGAATAAAAWTAGAAATCAATAACAAAAAG 724
Db
       369 GAAATTTGAAGAATCCATAAGTATGTGGAAATGAATCAAGGAACT----- 413
            725 ATACTCTGGAAAATNCACAAATACTTGGAAATTAAACAACATACTTCTAAATAACTCATG 784
Db
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Application/Control Number: 10/573,332 Art Unit: 1643

Qу	414	CAAGGGAAATTAGAAAATACTTTGAAATGAATGAAAATGAAAA	456
Db	785	$\tt GGTCAAAGAAATCAAAAGAGAAATTAAAAAAATTTTTGAAATAAAT$	844
Qy	457	CACAGCATACCAAAAC-TTATGAGATGCAGCTAAAATAGTGCTTACAGAGAAATTAATAG	515
Db	845	YACAACATATCAAAAATTTATGGGATGCAGCTAAAGCAGTGCTTAGAGGGAAATTTATAG	904
QУ	516	CTATTAATGCCTGTATTTTTTAAAAGAAGAAGATACCAAATTAAAAAAAA	575
Db	905	CATTAAATGCCTATATTAAAAAAGAAGAAGATCTCAAATCAATAACCTAAGTTTCCA	962
QУ	576	TTTAGAAAAAGAATAGTGAACCAAGCCCAAATCAAGCAGAAGGAAG	627
Db	963	CCTTAAGAAACTAGAAAAAGAAGAGCAAATTAAACCCAAAGTAAGCAGAAGAAAAGGAAAT	1022
Qy	628	AATAAAGATTAGAATGGAAAAATGAAATATGGAATTGGAAAAACTAGAGAAAAATTAA	687
Db	1023	AATAAAGATTAGAGCAGAAATAAATGAAATAGAAAACAGAAAAACAATAGAAAAAATCAA	1082
QУ	688	CAAACCCAAAAGTTGTTATATCAAAAAGATTGATAAGTTTGATAAACATTTAACTAGACT	747
Db	1083	TAAAACCAAAAGTTGGTTCTTTGAAAAGATAAAATTGACAAACCTTTAGCTAGACT	1138
Qу	748	TACCCTAATATCAAAACCACATACAGATATCACAAGAAAAGTACAGACCAATATCTCTCA	807
Db	1139	AAAAAAAAGAGAGAAGACACAAATTACTAATANATCAGAAATGAAAGAGAGAGAYATTACT	1198
Qy	808	TAAGACACATATAAGATAGA 827	
Db	1199	 ACAGATYCTACAGAAATAAA 1218	

Search results between Applicants' SEQ ID NO: 427 and Knoll sequence 148.

	ult No.	Score	Query Match	Length	DB	ID	Description
	1	1200	100.0	76360	16	US-10-669-920-427	Sequence 427, App
	2	1200	100.0	76360	19	US-10-573-332-427	Sequence 427, App
	3	939	78.2	75853	7	US-10-087-192-382	Sequence 382, App
С	4	506.6	42.2	665590	18	US-10-990-328-94275	Sequence 94275, A
С	5	484.6	40.4	161994	24	US-11-112-908-57	Sequence 57, Appl
С	6	484.6	40.4	303956	27	US-11-033-056A-36950	Sequence 36950, A
С	7	483	40.2	29142	10	US-10-741-600-17977	Sequence 17977, A
С	8	483	40.2	29142	17	US-10-796-280-12569	Sequence 12569, A
С	9	483	40.2	29142	36	US-12-287-505-17977	Sequence 17977, A
С	10	483	40.2	29142	36	US-12-337-905-17977	Sequence 17977, A
	11	473.8	39.5	130877	9	US-10-322-281-54	Sequence 54, Appl
	12	473.8	39.5	130877	16	US-10-539-228-54	Sequence 54, Appl
С	13	464.8	38.7	102139	18	US-10-990-328-95533	Sequence 95533, A
	14	464.8	38.7	439596	27	US-11-033-056A-37873	Sequence 37873, A
С	15	460.4	38.4	50498	17	US-10-105-299-9098	Sequence 9098, Ap
С	16	460.4	38.4	50498	18	US-10-868-184-6947	Sequence 6947, Ap
С	17	448.2	37.4	699032	18	US-10-990-328-96764	Sequence 96764, A
	18	446.2	37.2	330955	36	US-12-113-481-90	Sequence 90, Appl
	19	445	37.1	340000	24	US-11-102-978-3	Sequence 3, Appli
С	20	444.6	37.1	103000	27	US-11-033-056A-38751	Sequence 38751, A
С	21	441	36.8	2418	3	US-09-854-867-148	Sequence 148, App
C	22	441	36.8	2418	11	US-10-786-970A-148	Sequence 148, App

Application/Control Number: 10/573,332

Art Unit: 1643

7. Claims 33 and 35 are rejected under 35 U.S.C. 102(e) as being anticipated by Cargill et al./ U.S. Patent Application Publication number 2005/0026169 A1 (effective filing date April 30, 2003). Cargill discloses sequences that would selectively hybridize to Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, fragments thereof or their complements within in a sample, see sequence alignment information following instant rejection; page 3, sections 0023 and 0024; and page 14, section 0153. Cargill discloses sequences 17996 and 17977, which are the same as Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, respectively, corresponding hybridization probes, arrays, DNA chips and kits in which these components are contained, see abstract; page 14, sections 0154 and 0155; page 15, section 0162; and page 16, section 0173-page 18, section 0186.

Page 7

In absence of a clear definition of the term, electronic library as cited in the pending 112, 2nd paragraph rejection the art reads on claim 35. The Examiner regards the cited arrays and chips as the equivalent of Applicants' electronic library.

Search results between Applicants' SEQ ID NO: 427 and Cargill sequence 17977.

```
US-10-741-600-17977, rnpbm
; Sequence 17977, Application US/10741600; Publication No. US20050026169A1
; GENERAL INFORMATION:
 APPLICANT: CARGILL, Michele et al.
 TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
  TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF
 FILE REFERENCE: CL001499
 CURRENT APPLICATION NUMBER: US/10/741,600
 CURRENT FILING DATE: 2003-12-22
 NUMBER OF SEQ ID NOS: 73997
 SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17977
  LENGTH: 29142
  TYPE: DNA
  ORGANISM: Homo sapiens
US-10-741-600-17977
                  40.2%; Score 483; DB 10; Length 29142;
 Ouerv Match
 Best Local Similarity 73.0%;
                       0; Mismatches 215; Indels 76; Gaps 10;
 Matches 788; Conservative
Qу
       130 AAAAATACAAAAATTAGCTGGGTGTGGTGGTGCATGTCTGTAATCCCAGCTACTCAGGAG 189
          26298 AAAAATATTTTTTTTGGCCGAGTGCGGTGGCTCATGCCTGTAATCCCAGCTACTCAGGAG 26239
       190 GCTGAGGCTTGAACCTGG------GAGTCAGAGGTTGCAATGAGCCGAGATCGCAC 239
0.y
      26238 GCTGAGGCAGGAGATCGCTTGAACCTAGAAGCAGAGGTTGCAGTGAGCTGAGATCATGC 26179
Dh
       240 CACTGCACTCCAGCCTGGC-----GACAGAGCAAGACTCCTTGTCAAAA--AAAAAAA 290
QУ
          26178 CACGGCACTCCAGCCTGGCCGACAGAGCCGACGAAGGAGTTGGTCTCAACAGTGGCTTAA 26119
Db
       291 AAAATTCAGTAAACCATACTGTAAACAGATGTGCTGTCATTCAGGCTTAGTTATGCCATT 350
0v
          26118 \ \text{AATGTTCGGTAAACCATGCTGGGAACAGATGGGCTGTCATCCAGGCTTTGTTGTTCCATC} \ 26059
Db
       351 TACTGAACACAAGCAGAGTAGATTTAGCATAATTTCTAACAGCAATAGGATTTTTGAAAT 410
QУ
           26058 CACAGAGCACAGGCAGAATAGATTTAGCATCATTCTTAAGAGCCCTGGGATTTTCGGAAT 25999
Db
       411 GGTAAATGAACACTGGCTTCGACTTAAACTCACCAGCTGCATTACCTCCTAACAAGAGAG 470
Qу
          25998 GGTAAATGAGCATTGGCTTCAACTTAAAGTCGCCAGCTGCATTAGC-----CCTAACGAG 25944
Db
       471 AGTCAGCCTGTCCTTTGAAGCTCTGAAGCCAGACATTGACTT---ATCTGTAGCTAGAAA 527
Qy
          25943 AGTCAGCCTGTCCTTCGAAGCTTTGAAGCCAGGGATTGACTTCTCTTCTGTAGCTATGAA 25884
Db
       Q.y
          25883 AGTCCCAGGTGGCATCTTCTCCAATAGAAGGCTGTTTGTCTACATTGAAAGTCTGTTTT 25824
0.v
       579 -----AAATCTATGATCTTAGCTAGATCTTCTGGATGACTTGCTGCAGCT 623
                       Db
      25823 TGAGTGTAGCCACCTTCGTCAGTGAGATCAGCTAGATCTGATGGATAACTTACTGCAGCT 25764
       624 TCTACATCAGCACTTGCTTCACCTTGCACCTTTATTATATGGAGATGGCTTGTTTCC 683
QУ
          Db
      Qу
```

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25703 TTCAACCTCACAAAACAACCTCTGTTAGCTTCTGACTTTTCTTCTGCAACTTTCTCACCT 25644
Qу
       740 CCCTCAGCCTTCACAGAATTGAAGAGATTAGGGCATTGCTATGGATTAGGCTTTTGGCTT 799
          25643 CTCTCTGCCTTCAAAGGATTGAAGAGAGTGAGGGCCTTTCTCCGGACTAGGCTTTGGCTT 25584
Db
Qу
       800 AAGGGAATGTTATGGCTGGTTTGATCTTCTATCCAGATCACCCAAACTTTCCTCCATATC 859
          25583 AAAGGAATGTTGTGGCTGATTTGATCTTCTTTCCAGA----CTCAAACTTTCTCCGTATC 25528
Db
       860 AGCAATAATGCTGTTTTGCTTTTTTGTCATTCATATGTTCACTGGAGTAGCACTTTTAAT 919
Qу
          25527 AGCAATAATGCTATTCTGCTTTCATATCATTTGTGTGTCCACTAGAGCAGCACTTTTAAT 25468
Db
Ov
       920 T-----TTTTTTTTTTTTTTTCTTGATACAGAGTCTCGCTCTGTCACCCAGGC 963
                      25467 TTAATTTATCATTTTTATTTTTTTTTTTTTTTTTGAGACAGAATTTCGCTCTGTCACCCAGGT 25408
       964 TGGAGTGCAATGGCGTGGTCTCGGCTCACTGCAACCTCAGCCTCCTGGGTTCAAGTGGTT 1023
QУ
          Db
     25407 TGGAGTGCAGTGCTGCTCATGGTAACCTCCGCCTCCCCACTTCAAGCGATT 25348
Qу
      1024 CTCCCGCCTCAGCCTCCAGAGTAGCTGGGGCTACAGGGGCGCCACCACCACCACCCCAGC-CA 1082
          25347 CTCCTGCCTCAGCGTCCTAAGTAGTTCGGAGTACAGGCATGCACCACCACCACGCCAGCTAA 25288
Db
      1083 GTTTTTGTATTTTTAGTAGACACGGGGTTTCACTATGTTGGCCAGGCTGGACTCGAACT 1141
Qy
           25287 TTTTTTGTATTTTTAGTAGAGATGGGGTCTTGCCATGTTGGCCAGGCTGGTCTCGAACT 25229
```

rni. Search results between Applicants' SEQ ID NO: 4 and Cargill sequence 17996.

Result		Query				
No.	Score	Match I	Length D	B	ID	Description
1	289.6	24.1	3285	3	US-09-573-080A-425	Sequence 425, App
2	289.6	24.1		5	US-09-854-867-425	Sequence 425, App
3	245.8	20.5 2	278866	3	US-09-949-016-13922	Sequence 13922, A
4	245.8	20.5 2	278866	3	US-09-949-016-13923	Sequence 13923, A
5	245.8	20.5 2	278866	3	US-09-949-016-13924	Sequence 13924, A
6	245.8	20.5 2	278866	3	US-09-949-016-13925	Sequence 13925, A
7	245.8	20.5 2	278866	3	US-09-949-016-13926	Sequence 13926, A
8	245.8	20.5 2	278866	3	US-09-949-016-14699	Sequence 14699, A
9	245.8	20.5 2	278866	3	US-09-949-016-14700	Sequence 14700, A
10	245.8	20.5 2	278866	3	US-09-949-016-14701	Sequence 14701, A
11	245.8	20.5 2	278866	3	US-09-949-016-14702	Sequence 14702, A
12	245.8	20.5 2	278866	3	US-09-949-016-14703	Sequence 14703, A
13	232.8	19.4	999	10	US-10-301-480C-341210	Sequence 341210,
14	220.6	18.4 1	L48783	3	US-09-949-016-15729	Sequence 15729, A
15	215.2	17.9	998	10	US-10-301-480C-367650	Sequence 367650,
16	215.2	17.9	999	10	US-10-301-480C-267544	Sequence 267544,
17	215.2	17.9	999	10	US-10-301-480C-367649	Sequence 367649,
18	212.2	17.7	7678	3	US-09-573-080A-348	Sequence 348, App
19	212.2	17.7	7678	5	US-09-854-867-348	Sequence 348, App
c 20	211	.8 1	7.7 4	16	65 8 US-10-7 4 1-600	-17996 Sequence 17996, A

Art Unit: 1643

8. Claims 34 and 36 are rejected under 35 U.S.C. 102(e) as being anticipated by Venter et al./ U.S. Patent number 6,812,339 B1 (filed September 10, 2001). Venter discloses sequences that would selectively hybridize to Applicants' SEQ ID NO: 53 and SEQ ID NO: 622, fragments thereof or their complements within in a sample, see sequence alignment information following instant rejection; column 5, lines 32-47; and column 18, lines 24-63. Venter discloses sequences 340 and 3171, which are the same as Applicants' SEQ ID NO: 53 and SEQ ID NO: 622, respectively, corresponding hybridization probes, arrays, DNA chips, computer-based and data storage systems and kits in which these components are contained, see column 14, lines 39-54; and columns 15 and 16.

In absence of a clear definition of the term, electronic library as cited in the pending 112, 2nd paragraph rejection, the art reads on claim 35. The Examiner regards the cited arrays, chips and systems as the equivalent of Applicants' electronic library.

Search results between Applicants' SEQ ID NO: 53 and Venter sequence 340.

```
RESULT 3
US-09-949-016-340, rni
; Sequence 340, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
 APPLICANT: VENTER, J. Craig et al.
 TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
  TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
 FILE REFERENCE: CL001307
 CURRENT APPLICATION NUMBER: US/09/949,016
 CURRENT FILING DATE: 2000-04-14
 PRIOR APPLICATION NUMBER: 60/241,755
 PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
 PRIOR FILING DATE: 2000-10-03
 PRIOR APPLICATION NUMBER: 60/231,498
 PRIOR FILING DATE: 2000-09-08
 NUMBER OF SEQ ID NOS: 207012
  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 340
  LENGTH: 1145
  TYPE: DNA
  ORGANISM: Human
US-09-949-016-340
                   50.1%; Score 305.4; DB 3; Length 1145;
 Best Local Similarity 99.7%; Pred. No. 8.2e-75;
 Matches 306; Conservative 0; Mismatches
                                        Indels
Qу
        95 \ \ AGCAGGACAGGCTGCTTTGGTTTGTGACCTCCAGGCAGGACGGCCATCCTCTCCAGAATG \ 154
          Dh
        QУ
          Db
       215 ATGTGCTTCTCCTGCTTGAACCAGAAGAGCAATCTGTACTGCCTGAAGCCGACCATCTGC 274
Qу
          199 ATGTGCTTCTCCTGCTTGAACCAGAAGAGCAATCTGTACTGCCTGAAGCCGACCATCTGC 258
0.7
       275 TCCGACCAGGACAACTACTGCGTGACTGTCTGCTAGTGCCGGCATTGGGAATCTCGTG 334
          Db
       259 TCCGACCAGGACAACTACTGCGTGACTGTGTCTGCTAGTGCCGGCATTGGGAATCTCGTG 318
       335 ACATTTGGCCACAGCCTGAGCAAGACCTGTTCCCCGGCCTGCCCCATCCCAGAAGGCGTC 394
ΩУ
          Db
       319 ACATTTGGCCACAGCCTGAGCAAGACCTGTTCCCCGGCCTGCCCCATCCCAGAAGGCGTC 378
       395 AATGTGG 401
Q.y
          379 AATGTTG 385
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Page 12

Application/Control Number: 10/573,332

Art Unit: 1643

Search results between Applicants' SEQ ID NO: 622 and Venter sequences 3171 and 3172.

Resi		Score	Query Match	Length	DB	ID	Description	
	1	3270.8	79.0			US-10-648-593-132		
	2	3270.8	79.0				Sequence 132, App	
	3	2841.2	68.6	4268		US-09-954-556-3	Sequence 3, Appli	
	4	2525.8	61.0	3416		US-08-451-822A-15	Sequence 15, Appl	
	5	2525.8	61.0	3416	3		Sequence 15, Appl	
С		2425.6	58.6	4367	5	US-10-021-698A-3641	Sequence 3641, Ap	
	7	2413.8	58.3	3025	3	US-09-954-556-23	Sequence 23, Appl	
	8	2335	56.4	3080	3	US-09-954-556-25	Sequence 25, Appl	
	9	2269	54.8	3106	7	US-10-701-263-1	Sequence 1, Appli	
	10	2199.2	53.1	3244	3	US-09-954-556-24	Sequence 24, Appl	
С	11	2199.2	53.1	3244	5	US-10-021-698A-3626	Sequence 3626, Ap	
	12	1957	47.3	2310	2	US-08-471-570-9	Sequence 9, Appli	
	13	1949.6	47.1	2923	3	US-09-954-556-20	Sequence 20, Appl	
	14	1944.4	47.0	2826	3	US-09-954-556-21	Sequence 21, Appl	
	15	1943	46.9	2868	3	US-09-954-556-19	Sequence 19, Appl	
	16	1943	46.9	2941	3	US-09-954-556-18	Sequence 18, Appl	
	17	1834.4	44.3	2650	3	US-09-954-556-28	Sequence 28, Appl	
	18	1834.4	44.3	2676	2	US-08-471-570-7	Sequence 7, Appli	
	19	1688.4	40.8	3306	3	US-09-954-556-10	Sequence 10, Appl	
	20	1583	38.2		2		Sequence 3, Appli	
		1460.4	35.3		2	US-08-471-570-5	Sequence 5, Appli	
	22			32.2		079 3 US-09-949-016		'n
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	23			32.2		079	=	P
	24	1332.8	32.2	2079	3	US-09-949-016-3173	Sequence 3173, Ap	

9. Any inquiry concerning this communication or earlier communications from the Examiner should be directed to Alana M. Harris, Ph.D. whose telephone number is (571)272-0831. The Examiner works a flexible schedule, however she can normally be reached Monday through Saturday, 7:30 am to 6:30 pm with alternate Fridays off.

If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's supervisor, Larry R. Helms, Ph.D. can be reached on (571) 272-0832. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Art Unit: 1643

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Alana M. Harris, Ph.D. 26 February 2010 /Alana M. Harris, Ph.D./ Primary Examiner, Art Unit 1643